

Sequence Listing

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<120> Reagents and Methods Useful for Detecting Diseases of the Breast

<130> 6193.US.P1

<150> 08/971,772

<151> 17-Nov-1997

<160> 23

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 288

<212> DNA

<213> Homo sapiens

<400> 1

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aggcctggag	accagctccg	gtgggaagct	ggctggccat	cagaagaccg	tccccacggc	180
tcacctgact	tttgttattg	actgcaccca	cgggaagcag	ctctccctgg	cagcaaccgc	240
atcaccaccc	caagccccca	gtcccaatcg	agggttgtca	ccccacca		288

<210> 2

<211> 250

<212> DNA

<213> Homo sapiens

<400> 2

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gtgggaagct	ggctggccat	cagaagaccg	tccccacggc	tcacctgact	tttgttattg	120
actgcaccca	cgggaagcag	ctctccctgg	cagcaaccgc	atcaccaccc	caagccccca	180
gtcccaatcg	agggcttgct	acccaccaa	tgaagaccta	catcgtgttc	tgtggggaaa	240
actggcccca						250

<210> 3

<211> 256

<212> DNA

<213> Homo sapiens

<400> 3

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ccaggccagg	gccaccctgc	cgctctgcag	agggtctgtg	gcctcagctt	ccttcccagt	120
cagcccgtc	tgcccccagg	aggttcccg	ggctaagggg	aaaccctga	aggctgcgcc	180
tgtgaggtct	tcaacttggg	gaacagtcaa	ggactcactg	aaagccctct	cctcttgtgt	240
ctgtgggcag	gccgat					256

<210> 4

<211> 256

<212> DNA

<213> Homo sapiens

<400> 4

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ccagtaatat	ttgctgtatg	aatgaatgag	tctcttcatt	tgcagggtgac	ttatcctgcc	180
tctgccactc	gacggatgtt	tcagatgccc	cttagcggat	ctaattgatgt	ttccttggct	240
caagcacaaa	agactc					256

<210> 5

<211> 133

<212> DNA

<213> Homo sapiens

<400> 5

gctgttcaaa	atcatcttct	ttattttattg	gggttacttta	tttatttcagg	gtgggggtccc	60
tccaccccaa	aaataccagc	tccaggaaaa	ccatgggtatc	tccccagcac	tttgtagggc	120
ctggcatgtg	gaa					133

<210> 6

<211> 910

<212> DNA

<213> Homo sapiens

<400> 6

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aggcctggag	accagctccg	gtgggaagct	ggctggccat	cagaagaccg	tccccacggc	180
tcacctgact	tttgttattg	actgcaccca	cggaagcag	ctctccctgg	cagcaaccgc	240
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catcgtgttc	tgtggggaaa	actggcccca	tcttactcgg	gtgaccccca	tgggtggggg	360
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ggataagtca	cctgcacatg	aagagactca	ttcattcata	cagcaaata	tactggtaca	780
tcttccacat	gccaggccct	gcaaagtgtc	ggggagatac	catgggtttc	ctggagctgg	840
tatttttggg	gtggagggaa	cccaccctga	ataaataaag	taaccaata	aataaagaag	900
atgattttga						910

<210> 7

<211> 915

<212> DNA

<213> Homo sapiens

<400> 7

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aggcctggag	accagctccg	gtgggaagct	ggctggccat	cagaagaccg	tccccacggc	180
tcacctgact	tttgttattg	actgcaccca	cggaagcag	ctctccctgg	cagcaaccgc	240
atcaccaccc	caagccccca	gtcccaatcg	agggtctgtc	acccaccaa	tgaagacctt	300
catcgtgttc	tgtggggaaa	actggcccca	tctkactcgg	gtgaccccca	tgggtggggg	360
atgccttgcc	caggccaggg	ccaccctgcc	gctctgcaga	gggtctgtgg	cctcagcttc	420
cttcccagtc	agcccgtctc	gccccagga	gggtcccgag	gctaagggga	aaccctgtga	480
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gcaattccca	gggcctggcc	ctgcttcccc	agctaagcag	gagtcttttg	tgcttgagcc	660
aaggaaacat	cattagatcc	gctaaggggc	atctgaaaca	tccgtcgagt	ggcagaggca	720
ggataagtca	cctgcacatg	aagagactca	ttcattcata	cagcaaata	tactggtaca	780
tcttccacat	gccaggccct	gcaaagtgtc	ggggagatac	catgggtttc	ctggagctgg	840
tatttttggg	gtggagggaa	cccaccctga	ataaataaag	taaccaata	aataaagaag	900
atgattttga	acagc					915

<210> 8

<211> 68

<212> DNA
 <213> Artificial Sequence

 <220>
 <223> Restriction site

 <400> 8
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 cgggaatt 68

 <210> 9
 <211> 68
 <212> DNA
 <213> Artificial Sequence

 <400> 9
 aattaattcc cgggtcgacg agctcactag tcggcggccg ctctagagga tccaagctcg 60
 gaattccg 68

 <210> 10
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Universal primer

 <400> 10
 agcggataac aatttcacac agga 24

 <210> 11
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <400> 11
 tgtaaaacga cggccagt 18

 <210> 12
 <211> 20
 <212> DNA
 <213> Homo sapiens

 <400> 12
 cccaccaat gaagacctac 20

 <210> 13
 <211> 20
 <212> DNA
 <213> Homo sapiens

 <400> 13
 agaggagagg gctttcagt 20

 <210> 14
 <211> 20
 <212> DNA
 <213> Homo sapiens

 <400> 14
 cccacagaa cacgatgtag 20

 <210> 15
 <211> 22
 <212> DNA
 <213> Homo sapiens

<400> 15
ttgtcacccc accaatgaag ac

22

<210> 16
<211> 22
<212> DNA
<213> Homo sapiens

<400> 16
tggtatctcc ccagcacttt gc

22

<210> 17
<211> 188
<212> PRT
<213> Homo sapiens

<400> 17
Glu Trp Pro Arg Thr Ala Pro Leu Leu Pro Glu Leu Gly Arg Arg Arg
1 5 10 15
Ser Ser Arg Met Ala Pro Ser Glu Asp Pro Arg Asp Trp Arg Ala Asn
20 25 30
Leu Lys Gly Thr Ile Arg Glu Thr Gly Leu Glu Thr Ser Ser Gly Gly
35 40 45
Lys Leu Ala Gly His Gln Lys Thr Val Pro Thr Ala His Leu Thr Phe
50 55 60
Val Ile Asp Cys Thr His Gly Lys Gln Leu Ser Leu Ala Ala Thr Ala
65 70 75 80
Ser Pro Pro Gln Ala Pro Ser Pro Asn Arg Gly Leu Val Thr Pro Pro
85 90 95
Met Lys Thr Tyr Ile Val Phe Cys Gly Glu Asn Trp Pro His Leu Thr
100 105 110
Arg Val Thr Pro Met Gly Gly Gly Cys Leu Ala Gln Ala Arg Ala Thr
115 120 125
Leu Pro Leu Cys Arg Gly Ser Val Ala Ser Ala Ser Phe Pro Val Ser
130 135 140
Pro Leu Cys Pro Gln Glu Val Pro Glu Ala Lys Gly Lys Pro Val Lys
145 150 155 160
Ala Ala Pro Val Arg Ser Ser Thr Trp Gly Thr Val Lys Asp Ser Leu
165 170 175
Lys Ala Leu Ser Ser Cys Val Cys Gly Gln Ala Asp
180 185

<210> 18
<211> 21
<212> PRT
<213> Homo sapiens

<400> 18
Arg Ser Ser Arg Met Ala Pro Ser Glu Asp Pro Arg Asp Trp Arg Ala
1 5 10 15
Asn Leu Lys Gly Thr
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<210> 19
<211> 19
<212> PRT
<213> Homo sapiens

<400> 19
Met Gly Gly Gly Cys Leu Ala Gln Ala Arg Ala Thr Leu Pro Leu Cys
1 5 10 15
Arg Gly Ser

<210> 20
<211> 35
<212> PRT

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<213> Homo sapiens

<400> 20
Leu Cys Pro Gln Glu Val Pro Glu Ala Lys Gly Lys Pro Val Lys Ala
1      5      10      15
Ala Pro Val Arg Ser Ser Thr Trp Gly Thr Val Lys Asp Ser Leu Lys
20      25      30
Ala Leu Ser
35

<210> 21
<211> 19
<212> PRT
<213> Homo sapiens

<400> 21
Arg Glu Thr Gly Leu Glu Thr Ser Ser Gly Gly Lys Leu Ala Gly His
1      5      10      15
Gln Lys Thr

<210> 22
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Affinity purification system recognition site

<400> 22
Asp Tyr Lys Asp Asp Asp Asp Lys
1      5

<210> 23
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Affinity purification system recognition site

<400> 23
Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Met His Thr Glu His
1      5      10      15
His His His His His
20

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